

NA

(1643)

BATCH

1-12

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/323,597DATE: 02/01/2000
TIME: 17:11:34

Input Set: I323597.RAW

This Raw Listing contains the General Information
Section and up to first 5 pages.

ENTERED

P.S

1 <110> APPLICANT: Afar, Daniel E
2 Hubert, Rene S
3 Leong, Kahan
4 Raitano, Arthur B.
5 Saffran, Douglas C.
6 <120> TITLE OF INVENTION: NOVEL TUMOR ANTIGEN USEFUL IN DIAGNOSIS AND THERAPY OF
7 PROSTATE AND COLON CANCER
8 <130> FILE REFERENCE: 1703-007.US1
9 <140> CURRENT APPLICATION NUMBER: US/09/323,597
10 <141> CURRENT FILING DATE: 1999-06-01
11 <150> EARLIER APPLICATION NUMBER: 60/087,598
12 <151> EARLIER FILING DATE: 1998-06-01
13 <150> EARLIER APPLICATION NUMBER: 60/091,474
14 <151> EARLIER FILING DATE: 1998-06-29
15 <150> EARLIER APPLICATION NUMBER: 60/129,521
16 <151> EARLIER FILING DATE: 1999-04-14
17 <160> NUMBER OF SEQ ID NOS: 13
18 <170> SOFTWARE: PatentIn Ver. 2.0
19 <210> SEQ ID NO 1
20 <211> LENGTH: 1738
21 <212> TYPE: DNA
22 <213> ORGANISM: Homo sapiens
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26 aactcagggt caccaccagc tattggacct tactatgaaa accatggata ccaaccggaa 180
27 aaccctatc ccgcacagcc cactgtgtgc cccactgtct acgaggtgca tccggctcag 240
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29 gtcgtctgca cgcagcccaa atccccatcc gggacagtgt gcacctcaaa gactaagaaa 360
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34 cagaggaagt cctggcaccc tgtgtgccaa gacgactgga acgagaaacta cgggcggggc 660
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36 agcggatcca ccagctttat gaaactgaac acaagtgccg gcaatgtcga tatctataaa 780
37 aaactgtacc acagtgatgc ctgttcttca aaagcagtgg tttctttacg ctgtatagcc 840
38 tgcgggggtca acttgaactc aagccgccag agcaggattg tgggcgggca gagcgcgctc 900
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41 aatccatggc attggacggc atttgccggg attttgagac aatctttcat gttctatgga 1080
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44 gtgtgtctgc ccaacccagg catgatgctg cagccagaac agctctgctg gatttccggg 1260

RAW SEQUENCE LISTING
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TIME: 17:11:34

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47      atgatctgtg ccggcttcct gcaggggaac gtcgattctt gccaggggtga cagtggaggg 1440
48      cctctgggtca cttcgaagaa caatatctgg tggctgatag gggatacaag ctggggttct 1500
49      ggctgtgcc aagcttacag accaggagtg tacgggaatg tgatggtatt cacggactgg 1560
50      atttatcgac aaatgagggc agacggctaa tccacatggt cttcgtcctt gacgtcgttt 1620
51      tacaagaaaa caatggggct ggttttgcct ccccgatcat gatttactct tagagatgat 1680
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54 <211> LENGTH: 491

55 <212> TYPE: PRT

56 <213> ORGANISM: Homo sapiens

57 <400> SEQUENCE: 2

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61      20          25          30
62      Val Pro Thr Val Tyr Glu Val His Pro Ala Gln Tyr Tyr Pro Ser Pro
63      35          40          45
64      Val Pro Gln Tyr Ala Pro Arg Val Leu Thr Gln Ala Ser Asn Pro Val
65      50          55          60
66      Val Cys Thr Gln Pro Lys Ser Pro Ser Gly Thr Val Cys Thr Ser Lys
67      65          70          75          80
68      Thr Lys Lys Ala Leu Cys Ile Thr Leu Thr Leu Gly Thr Phe Leu Val
69      85          90          95
70      Gly Ala Ala Leu Ala Ala Gly Leu Leu Trp Lys Phe Met Gly Ser Lys
71      100          105          110
72      Cys Ser Asn Ser Gly Ile Glu Cys Asp Ser Ser Gly Thr Cys Ile Asn
73      115          120          125
74      Pro Ser Asn Trp Cys Asp Gly Val Ser His Cys Pro Gly Gly Glu Asp
75      130          135          140
76      Glu Asn Arg Cys Val Arg Leu Tyr Gly Pro Asn Phe Ile Leu Gln Val
77      145          150          155          160
78      Tyr Ser Ser Gln Arg Lys Ser Trp His Pro Val Cys Gln Asp Asp Trp
79      165          170          175
80      Asn Glu Asn Tyr Gly Arg Ala Ala Cys Arg Asp Met Gly Tyr Lys Asn
81      180          185          190
82      Asn Phe Tyr Ser Ser Gln Gly Ile Val Asp Asp Ser Gly Ser Thr Ser
83      195          200          205
84      Phe Met Lys Leu Asn Thr Ser Ala Gly Asn Val Asp Ile Tyr Lys Lys
85      210          215          220
86      Tyr His Ser Asp Ala Cys Ser Ser Lys Ala Val Val Ser Leu Arg Cys
87      225          230          235          240
88      Ile Ala Cys Gly Val Asn Leu Asn Ser Ser Arg Gln Ser Arg Ile Val
89      245          250          255
90      Gly Gly Glu Ser Ala Leu Pro Gly Ala Trp Pro Trp Gln Val Ser Leu
91      260          265          270
92      His Val Gln Asn Val His Val Cys Gly Gly Ser Ile Ile Thr Pro Glu
93      275          280          285
94      Trp Ile Val Thr Ala Ala His Cys Val Glu Lys Pro Leu Asn Asn Pro

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RAW SEQUENCE LISTING
PATENT APPLICATION US/09/323,597DATE: 02/01/2000
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98      Tyr Gly Ala Gly Tyr Gln Val Glu Lys Val Ile Ser His Pro Asn Tyr
99          325          330          335
100     Asp Ser Lys Thr Lys Asn Asn Asp Ile Ala Leu Met Lys Leu Gln Lys
101          340          345          350
102     Pro Leu Thr Phe Asn Asp Leu Val Lys Pro Val Cys Leu Pro Asn Pro
103          355          360          365
104     Gly Met Met Leu Gln Pro Glu Gln Leu Cys Trp Ile Ser Gly Trp Gly
105          370          375          380
106     Ala Thr Glu Glu Lys Gly Lys Thr Ser Glu Val Leu Asn Ala Ala Lys
107          385          390          395          400
108     Val Leu Leu Ile Glu Thr Gln Arg Cys Asn Ser Arg Tyr Val Tyr Asp
109          405          410          415
110     Asn Leu Ile Thr Pro Ala Met Ile Cys Ala Gly Phe Leu Gln Gly Asn
111          420          425          430
112     Val Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu Val Thr Ser Lys
113          435          440          445
114     Asn Asn Ile Trp Trp Leu Ile Gly Asp Thr Ser Trp Gly Ser Gly Cys
115          450          455          460
116     Ala Lys Ala Tyr Arg Pro Gly Val Tyr Gly Asn Val Met Val Phe Thr
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120 <210> SEQ ID NO 3

121 <211> LENGTH: 2479

122 <212> TYPE: DNA

123 <213> ORGANISM: Homo sapiens

124 <300> PUBLICATION INFORMATION:

125 <303> JOURNAL: Genomics

126 <304> VOLUME: 44

127 <306> PAGES: 309-320

128 <307> DATE: 1997

129 <400> SEQUENCE: 3

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132      cggaaaaccc ctatcccgca cagcccactg tggccccac tgtctacgag gtgcatccgg 180
133      ctccagtacta cccgtccccc gtgccccagt acgccccgag ggtcctgacg caggcttcca 240
134      accccgtcgt ctgcacgcag cccaaatccc catccgggac agtgtgcacc tcaaagacta 300
135      agaaagcact gtgcatcacc ttgaccctgg ggaccttcct cgtgggagct gcgctggccg 360
136      ctggccctact ctggaagttc atgggcagca agtgctccaa ctctgggata gagtgcgact 420
137      cctcaggtac ctgcatcaac ccctctaact ggtgtgatgg cgtgtcacac tgccccggcg 480
138      gggaggacga gaatcgggtg gttegcctct acggaccaa cttcatcctt cagatgtact 540
139      catctcagag gaatccttg caccctgtgt gccaaagcga ctggaacgag aactacgggc 600
140      gggcggcctg caggacatg ggctataaga ataattttta ctctagccaa ggaatagtgg 660
141      atgacagcgg atccaccagc tttatgaaac tgaacacaag tgccggcaat gtcgatatct 720
142      ataaaaaact gtaccacagt gatgcctgtt cttcaaaagc agtgggtttct ttacgctgtt 780
143      tagcctgcgg ggtcaacttg aactcaagcc gccagagcag gatcggtggc ggtgagagcg 840
144      cgctcccggg ggctggccc tggcaggtca gcctgcacgt ccagaacgct cacgtgtgcg 900

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RAW SEQUENCE LISTING
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147 atggagccgg ataccaagta caaaaagtga tttctcatcc aaattatgac tccaagacca 1080
148 agaacaatga cattgcgctg atgaagctgc agaagcctct gactttcaac gacctagtga 1140
149 aaccagtgtg tctgccccac ccaggcatga tgctgcagcc agaacagctc tgctggattt 1200
150 ccgggtgggg ggccaccgag gagaaaggga agacctcaga agtgctgaac gctgccaagg 1260
151 tgcttctcat tgagacacag agatgcaaca gcagatatgt ctatgacaac ctgatcacac 1320
152 cagccatgat ctgtgccggc ttcctgcagg ggaacgtcga ttcttgccag ggtgacagtg 1380
153 gagggcctct ggtcacttcg aacaacaata tctggtggct gataggggat acaagctggg 1440
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156 cgtttttacaa gaaaacaatg gggctggttt tgcttccccg tgcattgatt actcttagag 1620
157 atgattcaga ggtcacttca tttttattaa acagtgaact tgtctggctt tggcactctc 1680
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168 tcctagcacc ctggagagtg aatgccccct ggctccctggc agggcgccaa gtttggcacc 2340
169 atgtcggcct cttcaggcct gatagtcatt ggaaattgag gtccatgggg gaaatcaagg 2400
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174 <212> TYPE: PRT
175 <213> ORGANISM: Homo sapiens
176 <300> PUBLICATION INFORMATION:
177 <303> JOURNAL: Genomics
178 <304> VOLUME: 44
179 <306> PAGES: 309-320
180 <307> DATE: 1997
181 <400> SEQUENCE: 4

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182 Met Ala Leu Asn Ser Gly Ser Pro Pro Ala Ile Gly Pro Tyr Tyr Glu
183      1              5              10              15
184 Asn His Gly Tyr Gln Pro Glu Asn Pro Tyr Pro Ala Gln Pro Thr Val
185      20              25              30
186 Val Pro Thr Val Tyr Glu Val His Pro Ala Gln Tyr Tyr Pro Ser Pro
187      35              40              45
188 Val Pro Gln Tyr Ala Pro Arg Val Leu Thr Gln Ala Ser Asn Pro Val
189      50              55              60
190 Val Cys Thr Gln Pro Lys Ser Pro Ser Gly Thr Val Cys Thr Ser Lys
191      65              70              75              80
192 Thr Lys Lys Ala Leu Cys Ile Thr Leu Thr Leu Gly Thr Phe Leu Val
193      85              90              95
194 Gly Ala Ala Leu Ala Ala Gly Leu Leu Trp Lys Phe Met Gly Ser Lys

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RAW SEQUENCE LISTING
PATENT APPLICATION US/09/323,597DATE: 02/01/2000
TIME: 17:11:34

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199			130					135				140				
200	Glu	Asn	Arg	Cys	Val	Arg	Leu	Tyr	Gly	Pro	Asn	Phe	Ile	Leu	Gln	Met
201	145						150				155					160
202	Tyr	Ser	Ser	Gln	Arg	Lys	Ser	Trp	His	Pro	Val	Cys	Gln	Asp	Asp	Trp
203					165					170					175	
204	Asn	Glu	Asn	Tyr	Gly	Arg	Ala	Ala	Cys	Arg	Asp	Met	Gly	Tyr	Lys	Asn
205			180						185				190			
206	Asn	Phe	Tyr	Ser	Ser	Gln	Gly	Ile	Val	Asp	Asp	Ser	Gly	Ser	Thr	Ser
207			195					200				205				
208	Phe	Met	Lys	Leu	Asn	Thr	Ser	Ala	Gly	Asn	Val	Asp	Ile	Tyr	Lys	Lys
209			210					215				220				
210	Leu	Tyr	His	Ser	Asp	Ala	Cys	Ser	Ser	Lys	Ala	Val	Val	Ser	Leu	Arg
211	225					230					235					240
212	Cys	Leu	Ala	Cys	Gly	Val	Asn	Leu	Asn	Ser	Ser	Arg	Gln	Ser	Arg	Ile
213					245					250					255	
214	Val	Gly	Gly	Glu	Ser	Ala	Leu	Pro	Gly	Ala	Trp	Pro	Trp	Gln	Val	Ser
215			260						265				270			
216	Leu	His	Val	Gln	Asn	Val	His	Val	Cys	Gly	Gly	Ser	Ile	Ile	Thr	Pro
217			275					280				285				
218	Glu	Trp	Ile	Val	Thr	Ala	Ala	His	Cys	Val	Glu	Lys	Pro	Leu	Asn	Asn
219			290					295				300				
220	Pro	Trp	His	Trp	Thr	Ala	Phe	Ala	Gly	Ile	Leu	Arg	Gln	Ser	Phe	Met
221	305					310					315					320
222	Phe	Tyr	Gly	Ala	Gly	Tyr	Gln	Val	Gln	Lys	Val	Ile	Ser	His	Pro	Asn
223					325					330					335	
224	Tyr	Asp	Ser	Lys	Thr	Lys	Asn	Asn	Asp	Ile	Ala	Leu	Met	Lys	Leu	Gln
225			340						345				350			
226	Lys	Pro	Leu	Thr	Phe	Asn	Asp	Leu	Val	Lys	Pro	Val	Cys	Leu	Pro	Asn
227			355					360				365				
228	Pro	Gly	Met	Met	Leu	Gln	Pro	Glu	Gln	Leu	Cys	Trp	Ile	Ser	Gly	Trp
229			370					375				380				
230	Gly	Ala	Thr	Glu	Glu	Lys	Gly	Lys	Thr	Ser	Glu	Val	Leu	Asn	Ala	Ala
231	385					390					395					400
232	Lys	Val	Leu	Leu	Ile	Glu	Thr	Gln	Arg	Cys	Asn	Ser	Arg	Tyr	Val	Tyr
233					405					410					415	
234	Asp	Asn	Leu	Ile	Thr	Pro	Ala	Met	Ile	Cys	Ala	Gly	Phe	Leu	Gln	Gly
235			420						425				430			
236	Asn	Val	Asp	Ser	Cys	Gln	Gly	Asp	Ser	Gly	Gly	Pro	Leu	Val	Thr	Ser
237			435					440				445				
238	Asn	Asn	Asn	Ile	Trp	Trp	Leu	Ile	Gly	Asp	Thr	Ser	Trp	Gly	Ser	Gly
239			450					455				460				
240	Cys	Ala	Lys	Ala	Tyr	Arg	Pro	Gly	Val	Tyr	Gly	Asn	Val	Met	Val	Phe
241	465					470					475					480
242	Thr	Asp	Trp	Ile	Tyr	Arg	Gln	Met	Lys	Ala	Asn	Gly				
243					485					490						

Please Note:

<210> SEQ ID NO 5

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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VERIFICATION SUMMARY
PATENT APPLICATION US/09/323,597

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TIME: 17:11:34

Input Set: I323597.RAW

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253	W "N" or "Xaa" used: Feature required	agggaccagc ccttcattggg tggtagcgtg gtagtcac